

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q7Z3K3	POGZ_HUMAN	Homo sapiens	Pogo transposable element with ZNF domain	44.578178	S239;T240;S244;S246;T249;S251;T252;S256;T260;T262;T265;S266;S274;T282;S289;S292;S297;S300;T303;T310;S314;S333;S340;S343;S348;T361;S363;T429;S434;T699;T707;T719;S721;S1366;T1368	S333;S363;S425;T439;S445;T463;S856;S1338;S1364;S1367;T1368;S1373;S1374;T1378;S1392;T1394;S1396	32119511;34725712;35132862;31637018;29249144;23301498;32574038;22661428;30059200;20068230;35289036;33214551;30379171;34019948;35254053;28314751;32870666;37340703;29485866;27655845;28657654;27114449;34846842;30620550;31492838;29351928	MADTDLFMECEEEEELEPWQKISDVI EDSVVEDYNSVDKTTTVSVSQQPVS APVPIAAHASVAGHLSTSTTVSSSGA QNSDSTKKTTLVTLIANNNAGNPLVQ QGGQPLILTQNPAPGLGTMVTPVPL RPVQVMQANHVTSPPVASQPIFITT QGFVPRNVRPVQNAMNQVGIVLNV QQGQTVRPITLVPAPGTQFVKPTVGV PQVFSQMTVPVRPGSTMPVRPTTNTF TTVIPATLIRSTVPSQSQSQTKSTPS TSTTPTATQPTSLGQLAVQSPGQSN QTTNPKLAPSFPSPPAVSIASFVTVK RPGVTGENSNEVAKLVNLTNLTIPSL GQSPGPVVVSNSSAHGSQRTSGPE SSMKVTSIPVFDLQDGGKICPRC NAQFRVTEALRGHMCYCCPEMVEY QKKGKSLDSEPSVPSAAKPPSPEKTA PVASTPSSTPIPALSPPTKVPEPNENV GDAVQTKLIMLVDDFYGRDGGKVA QLTNFPKVATSFRCPHCTKRLKNNI RFMNMKHHVELDQQNGEVDGHT ICQHCRYQFSTPFQLQCHLENVHSP YESTTKCKICEWAFESEPLFLQHMK DTHKPGEMPYVCQVCQYRSSLYSEV DVHFRMIHEDTRHLLCPYCLKVFKN GNAFQQHYMRHQKRNVYHCNKCR LQFLFAKDKEHKLQHHKTFRKPQKQ LEGLKPGTKVTIRASRGQPRTPVSVS NDTPPSALQEAAPLTSMDPLPVFLY PPVQRSIQKRAVRKMSVMGRQTCLE CSFEIPDFPNHFPYVHCSLCRYSTC CSRAYANHMNNHVPKSPKYLALF KNSVSGIKLACTSCTFVTSVGDAMA KHLVFNPSHRSSSILPRGLTWIAHSR HGQTRDRVHDRVKNMYPPPSFPT NKAATVKSAGATPAEPEELLTPLAPA LPSPASTATPPPTPTHPQALALPLAT EGAECNLVDDQDEGSPVTQPELAS GGGGSGGVGKKEQLSVKCLRVLVFA LCCNTEQAAEHFRNPQRRIRRWLRR FQASQGENLEGKYLSFEAEKLAEW VLTQREQLPVNEETLFQKATKIGRS LEGGFKISYEWAVRFMLRHHLTPHA RRAVAHTLPKDVAENAGLFIDFVQR QIHNQDLPLSMIVAIDEISFLDTEVL SSDDRKENALQTVGTGEPWCDVVV AILADGTVLPTLVFYRGQMDQPANM PDSILLEAKESGYSDDDEIMELWSTRV WQKHTACQRSKGMVMDCHRTHL SEVLAMLASSTLPAVVPAGCSSKI QPLDVCIKRTVKNFLHKKWKEQARE MADTACDSVLLQLVVLVWLGVEVLG VIGDCPELVQRSFLVASVLPVPGPDGNI NSPTRNADMQEELIASLEEQKLSG EHSESSTPRRSPPEETIEPESLHQL FEGESETESFYGFEEADLDLMEI